

12. The method of any one of claims 7 to 11, wherein the epitopes comprising the polyepitope polypeptide are selected from epitopes of any one of the viruses of the group consisting of Epstein-Barr virus (EBV), hepatitis C virus (HCV), human immunodeficiency virus (HIV) and cytomegalovirus (CMV).

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13. A method of expressing a polypeptide in a suitable host, said method comprising, designing a polypeptide in accordance with the method of any one of claims 1 to 12, introducing a polynucleotide encoding said polypeptide into said host, such that said host is capable of expressing said polypeptide, and

10 culturing said bacterial host under conditions suitable for expression of said polypeptide.

14. A polypeptide designed in accordance with the method of any one of claims 1 to 5.

15. A polyepitope polypeptide designed in accordance with the method of any one of  
15 claims 1 to 12.

16. A polyepitope polypeptide comprising N epitopes, wherein N is any integer, said polyepitope polypeptide having the formula;

Triplet 1 - Triplet 2 - .... - Triplet N/3,

20 wherein each of said triplets comprises three linked epitopes selected by, identifying and ranking the relative hydrophobicity of each of the N epitopes, grouping the ranked N epitopes into three groups of substantially equivalent numbers, based upon the identified relative hydrophobicity of the N epitopes, to produce a first group comprising the epitopes of most relative hydrophobicity, a second group of epitopes of intermediate relative hydrophobicity, and a third group of epitopes of least relative hydrophobicity, and

25 selecting the epitopes for each of said triplets according to the following table:

Replaced By Article 19 Amendment  
SAC